RAW SEQUENCE LISTING PATENT APPLICATION US/08/823,980A

DATE: 10/28/97 / TIME: 09:11:18

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This Raw Listing contains the General Information Section and up to the first 5 pages.

	1		SEQUENCE LISTING	ERED
	2	(1)		CHED
	3 4	(1) G	General Information:	
	5 6	(i)	APPLICANT: Weiner, Amy J. Houghton, Michael	
	7 8 9	(ii)	TITLE OF INVENTION: Conserved Motif of Hepatitis C Viru E2/NS1 Region	s
	10 11 12	(iii)	NUMBER OF SEQUENCES: 134	
	13 14 15 16	(iv)	CORRESPONDENCE ADDRESS: (A) ADDRESSEE: Chiron Corporation (B) STREET: 4560 Horton Street (C) CITY: Emeryville	
•	17 18 19		(D) STATE: CA (E) COUNTRY: USA (F) ZIP: 94608-2916	
•	20 ²		COMPUTER READABLE FORM:	
. مد	22		(A) MEDIUM TYPE: Floppy disk	
	23 24 25	reak o	(B) COMPUTER: IBM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS (D) SOFTWARE: PatentIn Release #1.0, Version #1.25	
	26		, case — As althouses of	• •
	27	(V1)	CURRENT APPLICATION DATA:	
	28 29		(A) APPLICATION NUMBER: US 08/823,980 (B) FILING DATE: 25-MARCH-1997	
	30		(C) CLASSIFICATION:	
	31		(C) CHARRITICATION.	
	32	(viii)	ATTORNEY/AGENT INFORMATION:	
	33	, , , , ,	(A) NAME: Goldman, Kenneth M.	
	34		(B) REGISTRATION NUMBER: 34,174	
	35		(C) REFERENCE/DOCKET NUMBER: 0938.007	
	36			
	37	(ix)	TELECOMMUNICATION INFORMATION:	
	38		(A) TELEPHONE: 510-601-2719	
	39		(B) TELEFAX: 510-655-3542	
	40		. •	
	41	/2\ TNEO	DUARTON FOR ODO TO NO.1.	
	42 43	(Z) INFO	RMATION FOR SEQ ID NO:1:	
	44	/ i \	SEQUENCE CHARACTERISTICS:	
	45	(- /	(A) LENGTH: 6 amino acids	
	46		(B) TYPE: amino acid	

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```
(C) STRANDEDNESS: single
47
               (D) TOPOLOGY: linear
48
49
        (ii) MOLECULE TYPE: peptide
50
51
         (ix) FEATURE:
52
53
               (A) NAME/KEY: Xaa
               (B) LOCATION: 1
54
               (D) OTHER INFORMATION: /label= variable
55
                      /note= "Represents Ala, Arg, Asp, Gly, Lys, Ser,
56
57
58
         (ix) FEATURE:
59
               (A) NAME/KEY: Xaa
60
               (B) LOCATION: 2
61
               (D) OTHER INFORMATION: /label= variable
62
                      /note= "Represents Ile, Leu, Met, Phe or Trp"
63
64
         (ix) FEATURE:
65
               (A) NAME/KEY: Xaa
66
               (B) LOCATION: 3
67
               (D) OTHER INFORMATION: /label= variable
68
                      /note= "Represents Leu or Phe"
69
70
         (ix) FEATURE:
71
               (A) NAME/KEY: Xaa
72
73
               (B) LOCATION: 6
               (D) OTHER INFORMATION: /label= variable
74
                      /note= "Represents Ala or Gly"
75
76
77
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
78
79
          Xaa Xaa Xaa Xaa Xaa
80
                           5
81
          1
82
     (2) INFORMATION FOR SEQ ID NO:2:
83
84
          (i) SEQUENCE CHARACTERISTICS:
85
               (A) LENGTH: 32 amino acids
86
               (B) TYPE: amino acid
87
               (C) STRANDEDNESS: single
88
               (D) TOPOLOGY: linear
89
90
         (ii) MOLECULE TYPE: peptide
91
92
         (ix) FEATURE:
93
               (A) NAME/KEY: Xaa
94
               (B) LOCATEON: 1
95
               (C) OTHER INFORMATION: /label=Dpr
96
     /note=2,3-diaminopropionic is acetyl cysteine
97
98
99
         (ix) FEATURE:
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RAW SEQUENCE LISTING PATENT APPLICATION US/08/823,980A

The second section is

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100 101 102 103			 (A) NAME/KEY: Xaa (B) LOCATION: 2 (D) OTHER INFORMATION: /label=Ahe
104			
105		/ i \	SEQUENCE DESCRIPTION: SEQ ID NO:2:
106		(XI)	SEQUENCE DESCRIPTION. SEQ 15 No.1.
107		W	Xaa Glu Thr His Val Thr Gly Gly Ser Ala Gly His Thr Val Ser
108			
109		1	5 10 15
110		a1	Phe Val Ser Leu Leu Ala Pro Gly Ala Lys Gln Asn Val Gln Leu
111		GTA	20
112			20 25 30
113		T.V.T.O.1	DANGEON FOR CEO ID NO.2.
114	(2)	TWEO	RMATION FOR SEQ ID NO:3:
115			CROWNING CHARACTERICS.
116		(1)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 5 amino acids
117			
118			(B) TYPE: amino acid
119			(C) STRANDEDNESS: single
120			(D) TOPOLOGY: linear
121			VOLDOUED MUDE, montide
122		(11)	MOLECULE TYPE: peptide
123			
124			
125			THE PROPERTY OF THE PARTY OF TH
126		(X1)	SEQUENCE DESCRIPTION: SEQ ID NO:3:
127			
128			Ser Leu Leu Ala
129		1	5
130			
131	(2)	INFO	RMATION FOR SEQ ID NO:4:
132		_	
133		(i)	SEQUENCE CHARACTERISTICS:
134			(A) LENGTH: 7 amino acids
135			(B) TYPE: amino acid
136			(C) STRANDEDNESS: single
137			(D) TOPOLOGY: linear
138			Water Transport of the Control of th
139		(ii)	MOLECULE TYPE: peptide
140			
141			
142			THE PERSON AND THE WORLD
143		(Xi)	SEQUENCE DESCRIPTION: SEQ ID NO:4:
144			a a sia no dia sia
145			Leu Leu Ala Pro Gly Ala
146		1	5
147			DUALITAN TOP GEO TO NO. 5.
148	(2)	INFO	RMATION FOR SEQ ID NO:5:
149			CROWNING CHARACTERICAC.
150		(1)	SEQUENCE CHARACTERISTICS:
151			(A) LENGTH: 5 amino acids
167			CBA TEVERY AMERICACIO

205

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(C) STRANDEDNESS: single
153
               (D) TOPOLOGY: linear
154
155
         (ii) MOLECULE TYPE: peptide
156
157
158
159
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
160
161
          Leu Ala Pro Gly Ala
162
          1
                           5
163
164
     (2) INFORMATION FOR SEQ ID NO:6:
165
166
           (i) SEQUENCE CHARACTERISTICS:
167
                (A) LENGTH: 7 amino acids
168
                (B) TYPE: amino acid
169
                (C) STRANDEDNESS: single
170
                (D) TOPOLOGY: linear
171
172
          (ii) MOLECULE TYPE: peptide
173
174
175
176
          (ix) FEATURE:
                (A) NAME/KEY: Xaa
177
                (B) LOCATION: 7
178
                (D) OTHER INFORMATION: /label= variable
179
                       /note= "Represents Ala, Pro or Ser"
180
181
182
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
183
184
           Ser Leu Leu Xaa Xaa Gly Xaa
185
186
187
      (2) INFORMATION FOR SEQ ID NO:7:
188
189
           (i) SEQUENCE CHARACTERISTICS:
190
                (A) LENGTH: 7 amino acids
191
                (B) TYPE: amino acid
192
                (C) STRANDEDNESS: single
193
                (D) TOPOLOGY: linear
194
195
          (ii) MOLECULE TYPE: peptide
196
197
198
          (ix) FEATURE:
199
                (A) NAME/KEY: Xaa
200
                (B) LOCATION: 7
201
                (D) OTHER INFORMATION: /label= variable
202
                       /note= "Represents Ala, Pro or Ser"
203
204
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206 207		(Xi)	SEQUENCE	E DES	SCRIE	OITS	1: S1	EQ I	D NO	:7:					• •	
207		Ser	Leu Phe	Xaa	Xaa	Glv	Xaa									
209		1			5	2							÷		·	
210																
211																
212	(2)	INFO	RMATION I	OR S	SEQ]	D NO	:8:									
213	•															
214		(i)	SEQUENCE	CHA	ARAC'	PERIS	STICS	S:								
215			(A) LEI	VGTH:	31	amir	no ac	cids								
216			(B) TYE	PE: 8	amino	aci	ld									
217			(C) STE					Le							•	
218			(D) TOE	OLOC	3Y:]	linea	ar									
219							_									
220		(ii)	MOLECULE	TYE	PE: p	pepti	ide									
221																
222																
223																
224		/ ÷ \	CHOURNO	י בות	MDTI	эшт 🔿	. CI	20 TI	NIO.							
225 226		(XI)	SEQUENCE	r Des	CKI	1101	N: 21	20 T1) NO	:0:						
227		Yaa	Thr Xaa	Val	Пhr	G] v	G] v	Yaa	λla	α1ء	λra	Thr	Thr	Yaa	G] v	Yaa
228		1	IIII Auu	VUI	5	4 - y	O _T	nuu	niu	10	n- 9	****		nuu	15	nuu
229		-			•											
230		Xaa	Ser Leu	Phe	Xaa	Xaa	Glv	Xaa	Ser	Gln	Xaa	Ile	Gln	Leu	Ile	
231				20					25					30		
232																
233										•						
234	(2)	INFO	RMATION E	OR S	SEQ]	D NO):9:									
235																
236		(i)	SEQUENCE													
237			(A) LEN					cids								
238			(B) TYE													
239			(C) STE					це								
240 241			(D) TOE	OTO	L : Xi	Tue	ı L									
241		/ii\	MOLECULE	י ייטים) F. Y	ont i	do									
242		(11)	MOLECOLE	. 111	. r	Pepci	lue									
244																
245																
246																
247		(xi)	SEQUENCE	E DES	CRIE	OIT	l: SI	EQ II	ONO:	9:						
248		` '	_					_		,						
249		Ala	Thr Tyr	Ala	Thr	Gly	Ala	Ala	Gln	Gly	His	Ala	Thr	Asn	Ser	Phe
250		1	_		5	_				10					15	
251																
252		Val	Ser Leu	Phe	Ærg	Ser	Gly	Ala	Ser	Gln	Asn	Leu	Lys	Leu	Val	
253				20					25					30		
254				3		mac; m n										
255	(2)	INFO	RMATION E	OR S	SEQ]	D NO):10:	:								
256			9	<u></u>			.m = -									
257		(1)	SEQUENCE													
258			(A) LEŅ	IGTH:	1.5 - ا	amır	io ac	clds								

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SEQUENCE VERIFICATION REPORT PATENT APPLICATION *US/08/823,980A*

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